

GenCore version 6.3  
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:01:55 ; Search time 81 Seconds  
(without alignments)  
20559.273 Million cell

updates/sec

Title: US-09-300-482-1  
Perfect score: 233  
Sequence: 1  
gtttttgcagtttagtagaat.....ttggacaagggttattgttg 233

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 9875436 seqs, 3558593875 residues

Total number of hits satisfying chosen parameters: 19750872

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Issued\_Patents\_NA:\*  
1: /ABSS/Data/CRF/ptodata/1/ina/1\_COMB.seq:\*  
2: /ABSS/Data/CRF/ptodata/1/ina/5\_COMB.seq:\*  
3: /ABSS/Data/CRF/ptodata/1/ina/6A\_COMB.seq:\*  
4: /ABSS/Data/CRF/ptodata/1/ina/6B\_COMB.seq:\*  
5: /ABSS/Data/CRF/ptodata/1/ina/7A\_COMB.seq:\*  
6: /ABSS/Data/CRF/ptodata/1/ina/7B\_COMB.seq:\*  
7: /ABSS/Data/CRF/ptodata/1/ina/7C\_COMB.seq:\*  
8: /ABSS/Data/CRF/ptodata/1/ina/7D\_COMB.seq:\*  
9: /ABSS/Data/CRF/ptodata/1/ina/HA\_COMB.seq:\*  
10: /ABSS/Data/CRF/ptodata/1/ina/HB\_COMB.seq:\*  
11: /ABSS/Data/CRF/ptodata/1/ina/PCTUS\_COMB.seq:\*  
12: /ABSS/Data/CRF/ptodata/1/ina/PP\_COMB.seq:\*  
13: /ABSS/Data/CRF/ptodata/1/ina/RE\_COMB.seq:\*  
14: /ABSS/Data/CRF/ptodata/1/ina/backfiles1.seq:\*